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TECH CENTER 1600/2900
DATE: 10/17/2001
TIME: 15:51:31

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/641,892B

Input Set : A:\ES.txt
Output Set: N:\CRF3\10172001\I641892B.raw

corrected
version

ENTERED

3 <110> APPLICANT: IZUI, HIROSHI
4 MORIYA, MIKA
5 HIRANO, SEIKO
6 HARA, YOSHIHIKO
7 ITO, HISAO
8 MATSUI, KAZUHIKO
10 <120> TITLE OF INVENTION: METHOD FOR PRODUCING L-GLUTAMIC ACID BY FERMENTATION
ACCOMPANIED BY
11 PRECIPITATION
13 <130> FILE REFERENCE: 195942US0
15 <140> CURRENT APPLICATION NUMBER: 09/641,892B
16 <141> CURRENT FILING DATE: 2000-08-18
18 <150> PRIOR APPLICATION NUMBER: JP2000-78771
19 <151> PRIOR FILING DATE: 2000-03-21
21 <150> PRIOR APPLICATION NUMBER: JP11-234806
22 <151> PRIOR FILING DATE: 1999-08-20
24 <160> NUMBER OF SEQ ID NOS: 12
26 <170> SOFTWARE: PatentIn version 3.0
28 <210> SEQ ID NO: 1
29 <211> LENGTH: 4556
30 <212> TYPE: DNA
31 <213> ORGANISM: Enterobacter agglomerans
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34 <221> NAME/KEY: CDS
35 <222> LOCATION: (2)..(121)
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38 <221> NAME/KEY: CDS
39 <222> LOCATION: (322)..(3129)
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42 <221> NAME/KEY: CDS
43 <222> LOCATION: (3145)..(4368)
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46 <221> NAME/KEY: CDS
47 <222> LOCATION: (4437)..(4556)
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52 1 5 10 15
54 tgt cct aaa ggg cta aac ccg acg cgc gct atc ggc cac att aag tcg 97
55 Cys Pro Lys Gly Leu Asn Pro Thr Arg Ala Ile Gly His Ile Lys Ser
56 20 25 30
58 atg ctg ctg caa cgc agc gcg tag ttataccacc gggaacctca ggttcccgtt 151
59 Met Leu Leu Gln Arg Ser Ala
60 35
62 atttttacgga agcctctgta aacgcggtcc caaccacggtt tacaaagggtt cccttacggg 211
64 ccggggcgcgc gctgcgcaca gtgctcgat cgctgaactc actacggcaa accgcgaaa 271
66 cggcaacaaa tgaaacctca aaaaagcata acattgctta agggatcaca atg cag 327
67 Met Gln

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70 aac agc gcg atg aag ccc tgg ctg gac tcc tcc tgg ctg gcc ggc gcg      375
71 Asn Ser Ala Met Lys Pro Trp Leu Asp Ser Ser Trp Leu Ala Gly Ala
72                                     45                                     50                                     55
74 aat cag tct tac ata gag caa ctc tat gag gat ttc ctg acc gat cct      423
75 Asn Gln Ser Tyr Ile Glu Gln Leu Tyr Glu Asp Phe Leu Thr Asp Pro
76                                     60                                     65                                     70
78 gac tct gtg gat gca gtg tgg cgc tcg atg ttc caa cag tta cca ggc      471
79 Asp Ser Val Asp Ala Val Trp Arg Ser Met Phe Gln Gln Leu Pro Gly
80                                     75                                     80                                     85
82 acg gga gtg aaa cct gag cag ttc cac tcc gca act cgc gaa tat ttc      519
83 Thr Gly Val Lys Pro Glu Gln Phe His Ser Ala Thr Arg Glu Tyr Phe
84 90                                     95                                     100                                     105
86 cgt cgc ctg gcg aaa gac gca tct cgt tac acc tcc tca gtt acc gat      567
87 Arg Arg Leu Ala Lys Asp Ala Ser Arg Tyr Thr Ser Ser Val Thr Asp
88                                     110                                     115                                     120
90 ccg gca acc aac tcc aaa caa gtg aaa gtg ctg cag ctg att aac gcg      615
91 Pro Ala Thr Asn Ser Lys Gln Val Lys Val Leu Gln Leu Ile Asn Ala
92                                     125                                     130                                     135
94 ttt cgt ttc cgc gga cat cag gaa gca aat ctc gat ccg ctt ggc ctg      663
95 Phe Arg Phe Arg Gly His Gln Glu Ala Asn Leu Asp Pro Leu Gly Leu
96                                     140                                     145                                     150
98 tgg aaa cag gac cgc gtt gcc gat ctc gat cct gcc ttt cac gat ctg      711
99 Trp Lys Gln Asp Arg Val Ala Asp Leu Asp Pro Ala Phe His Asp Leu
100                                     155                                     160                                     165
102 acc gac gcc gat ttt cag gaa agc ttt aac gta ggt tct ttt gcc att      759
103 Thr Asp Ala Asp Phe Gln Glu Ser Phe Asn Val Gly Ser Phe Ala Ile
104 170                                     175                                     180                                     185
106 ggc aaa gaa acc atg aag ctg gcc gat ctg ttc gac gcg ctg aag cag      807
107 Gly Lys Glu Thr Met Lys Leu Ala Asp Leu Phe Asp Ala Leu Lys Gln
108                                     190                                     195                                     200
110 acc tac tgt ggc tcg att ggt gca gag tat atg cac atc aat aac acc      855
111 Thr Tyr Cys Gly Ser Ile Gly Ala Glu Tyr Met His Ile Asn Asn Thr
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114 gaa gag aaa cgc tgg atc cag cag cgt atc gaa tcc ggt gcg agc cag      903
115 Glu Glu Lys Arg Trp Ile Gln Gln Arg Ile Glu Ser Gly Ala Ser Gln
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118 acg tca ttc agt ggc gaa gag aaa aaa ggt ttc ctg aaa gag ctg acc      951
119 Thr Ser Phe Ser Gly Glu Glu Lys Lys Gly Phe Leu Lys Glu Leu Thr
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123 Ala Ala Glu Gly Leu Glu Lys Tyr Leu Gly Ala Lys Phe Pro Gly Ala
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126 aaa cgt ttc tcg ctg gaa ggc ggt gat gcg ctg gtg ccg atg ctg cgc      1047
127 Lys Arg Phe Ser Leu Glu Gly Gly Asp Ala Leu Val Pro Met Leu Arg
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130 gag atg att cgt cat gcg ggc aaa agc ggc aca cgt gaa gtg gta ctg      1095
131 Glu Met Ile Arg His Ala Gly Lys Ser Gly Thr Arg Glu Val Val Leu
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136			300					305					310				
138	aaa	aag	cca	cag	gat	ctg	ttc	gac	gaa	ttc	tcc	ggt	aaa	cac	aaa	gag	1191
139	Lys	Lys	Pro	Gln	Asp	Leu	Phe	Asp	Glu	Phe	Ser	Gly	Lys	His	Lys	Glu	
140		315					320					325					
142	cat	ctg	ggc	acc	ggt	gat	gtg	aag	tat	cac	atg	ggc	ttc	tct	tcg	gat	1239
143	His	Leu	Gly	Thr	Gly	Asp	Val	Lys	Tyr	His	Met	Gly	Phe	Ser	Ser	Asp	
144	330					335				340					345		
146	att	gaa	acc	gaa	ggt	ggt	ctg	gtg	cat	ctg	gcg	ctg	gcg	ttt	aac	ccg	1287
147	Ile	Glu	Thr	Glu	Gly	Gly	Leu	Val	His	Leu	Ala	Leu	Ala	Phe	Asn	Pro	
148				350					355						360		
150	tct	cac	ctg	gaa	att	gtc	agc	ccg	gtg	gtc	atg	gga	tcg	gta	cgt	gca	1335
151	Ser	His	Leu	Glu	Ile	Val	Ser	Pro	Val	Val	Met	Gly	Ser	Val	Arg	Ala	
152			365					370					375				
154	cgt	ctc	gat	cgt	ctg	gcc	gaa	ccg	gtc	agc	aat	aaa	gtg	ttg	cct	atc	1383
155	Arg	Leu	Asp	Arg	Leu	Ala	Glu	Pro	Val	Ser	Asn	Lys	Val	Leu	Pro	Ile	
156		380					385					390					
158	acc	att	cac	ggt	gat	gcg	gcg	gtg	att	ggt	cag	ggc	gtg	gtt	cag	gaa	1431
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162	acc	ctg	aac	atg	tct	cag	gcg	cgc	ggc	tac	gaa	gtg	ggc	ggc	acg	gta	1479
163	Thr	Leu	Asn	Met	Ser	Gln	Ala	Arg	Gly	Tyr	Glu	Val	Gly	Gly	Thr	Val	
164	410					415			420						425		
166	cgt	atc	gtc	att	aac	aac	cag	gtt	ggt	ttt	acc	acc	tcc	aac	ccg	aaa	1527
167	Arg	Ile	Val	Ile	Asn	Asn	Gln	Val	Gly	Phe	Thr	Thr	Ser	Asn	Pro	Lys	
168				430				435							440		
170	gat	gcg	cgt	tca	acc	ccg	tac	tgt	act	gac	atc	ggc	aag	atg	gtg	ctg	1575
171	Asp	Ala	Arg	Ser	Thr	Pro	Tyr	Cys	Thr	Asp	Ile	Gly	Lys	Met	Val	Leu	
172				445				450							455		
174	gca	ccg	att	ttc	cac	gtc	aat	gct	gac	gat	ccg	gaa	gcg	gtg	gcc	ttt	1623
175	Ala	Pro	Ile	Phe	His	Val	Asn	Ala	Asp	Asp	Pro	Glu	Ala	Val	Ala	Phe	
176		460					465					470					
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183	Phe	Ile	Asp	Leu	Val	Cys	Tyr	Arg	Arg	His	Gly	His	Asn	Glu	Ala	Asp	
184	490				495					500					505		
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187	Glu	Pro	Ser	Ala	Thr	Gln	Pro	Leu	Met	Tyr	Gln	Lys	Ile	Lys	Lys	His	
188				510				515							520		
190	ccg	acg	ccg	cgt	aaa	att	tac	gcc	gat	cgt	ctg	gaa	ggc	gaa	ggt	gtc	1815
191	Pro	Thr	Pro	Arg	Lys	Ile	Tyr	Ala	Asp	Arg	Leu	Glu	Gly	Glu	Gly	Val	
192				525				530							535		
194	gcg	tcg	cag	gaa	gat	gcc	acc	gag	atg	gtg	aac	ctg	tac	cgc	gat	gcg	1863
195	Ala	Ser	Gln	Glu	Asp	Ala	Thr	Glu	Met	Val	Asn	Leu	Tyr	Arg	Asp	Ala	
196		540					545					550					
198	ctc	gat	gcg	ggc	gaa	tgc	gtg	gtg	ccg	gaa	tgg	cgt	ccg	atg	agc	ctg	1911

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203	His	Ser	Phe	Thr	Trp	Ser	Pro	Tyr	Leu	Asn	His	Glu	Trp	Asp	Glu	Pro	
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206	tat	ccg	gca	cag	gtt	gac	atg	aaa	cgc	ctg	aag	gaa	ctg	gca	ttg	cgt	2007
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208					590					595					600		
210	atc	agc	cag	gtc	cct	gag	cag	att	gaa	gtg	cag	tcg	cgc	gtg	gcc	aag	2055
211	Ile	Ser	Gln	Val	Pro	Glu	Gln	Ile	Glu	Val	Gln	Ser	Arg	Val	Ala	Lys	
212				605					610					615			
214	atc	tat	aac	gat	cgc	aag	ctg	atg	gcc	gaa	ggc	gag	aaa	gcg	ttc	gac	2103
215	Ile	Tyr	Asn	Asp	Arg	Lys	Leu	Met	Ala	Glu	Gly	Glu	Lys	Ala	Phe	Asp	
216			620					625					630				
218	tgg	ggc	ggt	gcc	gag	aat	ctg	gcg	tac	gcc	acg	ctg	gtg	gat	gaa	ggt	2151
219	Trp	Gly	Gly	Ala	Glu	Asn	Leu	Ala	Tyr	Ala	Thr	Leu	Val	Asp	Glu	Gly	
220			635				640					645					
222	att	ccg	gtt	cgc	ctc	tcg	ggt	gaa	gac	tcc	ggt	cgt	gga	acc	ttc	ttc	2199
223	Ile	Pro	Val	Arg	Leu	Ser	Gly	Glu	Asp	Ser	Gly	Arg	Gly	Thr	Phe	Phe	
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226	cat	cgc	cac	gcg	gtc	gtg	cac	aac	cag	gct	aac	ggt	tca	acc	tat	acg	2247
227	His	Arg	His	Ala	Val	Val	His	Asn	Gln	Ala	Asn	Gly	Ser	Thr	Tyr	Thr	
228					670					675					680		
230	ccg	ctg	cac	cat	att	cat	aac	agc	cag	ggc	gag	ttc	aaa	gtc	tgg	gat	2295
231	Pro	Leu	His	His	Ile	His	Asn	Ser	Gln	Gly	Glu	Phe	Lys	Val	Trp	Asp	
232					685					690					695		
234	tcg	gtg	ctg	tct	gaa	gaa	gcg	gtg	ctg	gcg	ttt	gaa	tac	ggt	tac	gcc	2343
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250	ggc	tac	gaa	ggt	cag	gga	ccg	gaa	cac	tcc	tct	gcc	cgt	ctg	gaa	cgc	2535
251	Gly	Tyr	Glu	Gly	Gln	Gly	Pro	Glu	His	Ser	Ser	Ala	Arg	Leu	Glu	Arg	
252					765					770					775		
254	tat	ctg	caa	ctt	tgc	gcc	gag	cag	aac	atg	cag	gtt	tgc	gtc	ccg	tcg	2583
255	Tyr	Leu	Gln	Leu	Cys	Ala	Glu	Gln	Asn	Met	Gln	Val	Cys	Val	Pro	Ser	
256					780					785					790		
258	acg	ccg	gct	cag	gtg	tat	cac	atg	ctg	cgc	cgt	cag	gcg	ctg	cgc	ggg	2631
259	Thr	Pro	Ala	Gln	Val	Tyr	His	Met	Leu	Arg	Arg	Gln	Ala	Leu	Arg	Gly	
260					795				800						805		
262	atg	cgc	cgt	ccg	ctg	gtg	gtg	atg	tcg	ccg	aag	tcg	ctg	tta	cgc	cat	2679
263	Met	Arg	Arg	Pro	Leu	Val	Val	Met	Ser	Pro	Lys	Ser	Leu	Leu	Arg	His	

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270	ccg	gcc	att	ggt	gag	atc	gac	gat
271	Pro	Ala	Ile	Gly	Glu	Ile	Asp	Asp
272			845				850	
274	gtc	gtg	ctg	tgc	tcc	ggt	aag	ggt
275	Val	Val	Leu	Cys	Ser	Gly	Lys	Val
276			860				865	
278	cgt	aaa	gac	gag	aaa	acc	gat	ggt
279	Arg	Lys	Asp	Glu	Lys	Thr	Asp	Val
280		875				880		885
282	tac	ccg	ttc	ccg	cat	cag	gcg	gta
283	Tyr	Pro	Phe	Pro	His	Gln	Ala	Val
284	890				895		900	
286	cac	gta	cag	gac	ttt	gtc	tgg	tgc
287	His	Val	Gln	Asp	Phe	Val	Trp	Cys
288				910			915	
290	gcc	tgg	tac	tgt	agc	cag	cat	cat
291	Ala	Trp	Tyr	Cys	Ser	Gln	His	His
292			925				930	
294	gcc	acc	ctg	cgt	tat	gca	ggt	cgc
295	Ala	Thr	Leu	Arg	Tyr	Ala	Gly	Arg
296			940				945	
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300		955				960		965
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303	Ala	Leu	Asn	Val	Asn			Met
304	970						975	
306	ctc	ggt	ccc	gac	ctg	cct	gaa	tcg
307	Leu	Val	Pro	Asp	Leu	Pro	Glu	Ser
308				985			990	
310	tgg	cac	aag	aaa	cca	ggc	gat	gca
311	Trp	His	Lys	Lys	Pro	Gly	Asp	Ala
312			1000				1005	
314	gtc	gaa	att	gaa	act	gac	aaa	gtc
315	Val	Glu	Ile	Glu	Thr	Asp	Lys	Val
316			1015				1020	
318	gcc	gat	ggc	gtg	ctg	gaa	gcc	gtg
319	Ala	Asp	Gly	Val	Leu	Glu	Ala	Val
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